

V. Partner

1643

CRF Err Corrected by the STIC Systems Branch

Serial Number: 097545, 199C

CRF Processing Date: 8/11
Edited by: _____
Verified by: _____ (STIC sta

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TECH CENTER 1600/2900

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number input by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID.NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Corrected misaligned amino acid number in
sequence #16.

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/545,199C

DATE: 07/19/2001

TIME: 10:36:34

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\07192001\I545199C.raw

see p. 5

3 <110> APPLICANT: Lowery E., David
 4 Fuller E., Troy
 5 Kennedy J., Michael
 7 <120> TITLE OF INVENTION: Anti-Bacterial Vaccine Compositions
 9 <130> FILE REFERENCE: 28341/6227.1
 11 <140> CURRENT APPLICATION NUMBER: 09/545,199C
 12 <141> CURRENT FILING DATE: 2000-04-06
 14 <150> PRIOR APPLICATION NUMBER: 60/153,453
 15 <151> PRIOR FILING DATE: 1999-09-10
 17 <150> PRIOR APPLICATION NUMBER: 60/128,689
 18 <151> PRIOR FILING DATE: 1999-04-09
 20 <160> NUMBER OF SEQ ID NOS: 169
 22 <170> SOFTWARE: PatentIn Ver. 2.0
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 51 aacaatagtt ttaaacaata ttcttccatt ttttataagt aagtacttaa atataaagca 180
 53 ttttcataaa tatcaataaa ggattagtt atg gca gca gag ctt aca aca gcg 233
 54 Met Ala Ala Glu Leu Thr Thr Ala
 55 1 5
 59 gga tat att ggg cac cat tta gca ttc ttg aaa aca ggg gat tct ttc 281
 60 Gly Tyr Ile Gly His His Leu Ala Phe Leu Lys Thr Gly Asp Ser Phe
 61 10 15 20
 64 tgg cat gtt cat tta gat acc ctt cta ttt tca att att tca ggt gca 329
 65 Trp His Val His Leu Asp Thr Leu Leu Phe Ser Ile Ile Ser Gly Ala
 66 25 30 35 40
 68 att ttt ctt ttt gtt ttt tca aaa gtt gca aaa aaa gca acg ccg ggt 377
 69 Ile Phe Leu Phe Val Phe Ser Lys Val Ala Lys Lys Ala Thr Pro Gly
 70 45 50 55
 72 gtg cct agc aag atg caa tgt ttt gtt gag ata atg gtt gat tgg att 425

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Input Set : A:\Cpg.pto

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74          60          65          70
76 gat ggg atc gta aaa gaa aat ttc cat ggt cct cgt cat gct gtt gga 473
77 Asp Gly Ile Val Lys Glu Asn Phe His Gly Pro Arg His Ala Val Gly
78          75          80          85
80 cca tta gca tta act att ttc tgc tgg gta ttc att atg aat gct atc 521
81 Pro Leu Ala Leu Thr Ile Phe Cys Trp Val Phe Ile Met Asn Ala Ile
82          90          95          100
84 gat ttg atc cca gta gat ttc cta cct caa tta gcc cat tta ttt ggt 569
85 Asp Leu Ile Pro Val Asp Phe Leu Pro Gln Leu Ala His Leu Phe Gly
86 105          110          115          120
88 att gaa tac tta aga gct gtt cca aca gca gat atc agt gga aca tta 617
89 Ile Glu Tyr Leu Arg Ala Val Pro Thr Ala Asp Ile Ser Gly Thr Leu
90          125          130          135
92 ggc tta tca att ggt gtc ttc ttc tta att att ttc tat aca atc aaa 665
93 Gly Leu Ser Ile Gly Val Phe Phe Leu Ile Ile Phe Tyr Thr Ile Lys
94          140          145          150
96 tca aaa ggt atg agt ggc ttt gtt aaa gaa tat acg ctt cat cct ttt 713
97 Ser Lys Gly Met Ser Gly Phe Val Lys Glu Tyr Thr Leu His Pro Phe
98          155          160          165
100 aat cat cct ttg tta att ccg gtt aac tta gcg ctt gaa tca gtc aca 761
101 Asn His Pro Leu Leu Ile Pro Val Asn Leu Ala Leu Glu Ser Val Thr
102          170          175          180
104 tta tta gca aaa cct gtt tct ttg gcg ttc cgt ctt ttc ggg aat atg 809
105 Leu Leu Ala Lys Pro Val Ser Leu Ala Phe Arg Leu Phe Gly Asn Met
106 185          190          195          200
108 tat gca ggt gaa ctt atc ttt att ctt att gca gtg atg tac atg gca 857
109 Tyr Ala Gly Glu Leu Ile Phe Ile Leu Ile Ala Val Met Tyr Met Ala
110          205          210          215
112 aat aat ttt gca ctt aat tca atg ggt att ttc atg cat ttg gct tgg 905
113 Asn Asn Phe Ala Leu Asn Ser Met Gly Ile Phe Met His Leu Ala Trp
114          220          225          230
117 gct att ttc cat att ctt gtg att acc tta caa gca ttt att ttt atg 953
118 Ala Ile Phe His Ile Leu Val Ile Thr Leu Gln Ala Phe Ile Phe Met
119          235          240          245
121 atg ctt aca gtg gtt tat ttg agt atg ggt tat aac aaa gca gaa cac 1001
122 Met Leu Thr Val Val Tyr Leu Ser Met Gly Tyr Asn Lys Ala Glu His
123          250          255          260
125 taatttttta taaacaaaac cagaccttgg gtctaaattt caatcttatg gagaacatta 1061
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137 1 5 10 15
139 Phe Leu Lys Thr Gly Asp Ser Phe Trp His Val His Leu Asp Thr Leu
140 20 25 30

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143          35          40          45
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146          50          55          60
148 Val Glu Ile Met Val Asp Trp Ile Asp Gly Ile Val Lys Glu Asn Phe
149        65          70          75          80
151 His Gly Pro Arg His Ala Val Gly Pro Leu Ala Leu Thr Ile Phe Cys
152          85          90          95
154 Trp Val Phe Ile Met Asn Ala Ile Asp Leu Ile Pro Val Asp Phe Leu
155          100         105         110
157 Pro Gln Leu Ala His Leu Phe Gly Ile Glu Tyr Leu Arg Ala Val Pro
158          115         120         125
160 Thr Ala Asp Ile Ser Gly Thr Leu Gly Leu Ser Ile Gly Val Phe Phe
161          130         135         140
163 Leu Ile Ile Phe Tyr Thr Ile Lys Ser Lys Gly Met Ser Gly Phe Val
164 145          150         155         160
166 Lys Glu Tyr Thr Leu His Pro Phe Asn His Pro Leu Leu Ile Pro Val
167          165         170         175
169 Asn Leu Ala Leu Glu Ser Val Thr Leu Leu Ala Lys Pro Val Ser Leu
170          180         185         190
172 Ala Phe Arg Leu Phe Gly Asn Met Tyr Ala Gly Glu Leu Ile Phe Ile
173          195         200         205
175 Leu Ile Ala Val Met Tyr Met Ala Asn Asn Phe Ala Leu Asn Ser Met
176          210         215         220
178 Gly Ile Phe Met His Leu Ala Trp Ala Ile Phe His Ile Leu Val Ile
179 225          230         235         240
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195 <222> LOCATION: (364)..(1230)
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198 <223> OTHER INFORMATION: atpG
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205 acaacaagca ttagtggtat tcgcagtaga gtttggttac ttagaagaag tggacttaga 180
207 tcgtattggt tcatttgaat cagcactttt agagtatgct aaccataact atgctgattt 240
209 tatgctgtag ttaacccaat ctggcaatta caatgatgaa attaaagagt cattaataagg 300
211 cattttggat agcttcaaag caaacagtgc gtggttaagt aacactttta atggagagac 360
213 aaa atg gca ggt gct aaa gag ata aga acc aaa atc gcg agt gta aaa 408
214 Met Ala Gly Ala Lys Glu Ile Arg Thr Lys Ile Ala Ser Val Lys
215      1          5          10          15

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Input Set : A:\Cpg.pto

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218 Ser Thr Gln Lys Ile Thr Lys Ala Met Glu Met Val Ala Ala Ser Lys
219                20                25                30
221 atg cgt aaa acg caa gaa cgc atg tct tct tca cgc cct tat tca gaa 504
222 Met Arg Lys Thr Gln Glu Arg Met Ser Ser Ser Arg Pro Tyr Ser Glu
223                35                40                45
225 aca ata cgt aac gtg att agc cac gtt tcc aaa gca acg att ggt tac 552
226 Thr Ile Arg Asn Val Ile Ser His Val Ser Lys Ala Thr Ile Gly Tyr
227                50                55                60
229 aag cat cca ttt tta gtg gat cgc gaa gta aaa aaa gtg ggc atg att 600
230 Lys His Pro Phe Leu Val Asp Arg Glu Val Lys Lys Val Gly Met Ile
231                65                70                75
233 gtt gtg tcc aca gat cgt ggt ctt tgt ggt ggc tta aac gtg aac ttg 648
234 Val Val Ser Thr Asp Arg Gly Leu Cys Gly Gly Leu Asn Val Asn Leu
235                80                85                90                95
237 ttt aaa act gta tta aat gaa atg aaa gaa tgg aaa gaa aaa gat gtt 696
238 Phe Lys Thr Val Leu Asn Glu Met Lys Glu Trp Lys Glu Lys Asp Val
239                100                105                110
241 tcc gtt caa ttg agt tta atc ggt tct aaa tct atc aac ttt ttc caa 744
242 Ser Val Gln Leu Ser Leu Ile Gly Ser Lys Ser Ile Asn Phe Phe Gln
243                115                120                125
245 tct ttg gga att aaa att tta acc caa gat tca ggt att ggt gat act 792
246 Ser Leu Gly Ile Lys Ile Leu Thr Gln Asp Ser Gly Ile Gly Asp Thr
247                130                135                140
249 ccc tct gtt gag cag tta att ggt tca gtc aat tct atg att gat gct 840
250 Pro Ser Val Glu Gln Leu Ile Gly Ser Val Asn Ser Met Ile Asp Ala
251                145                150                155
253 tat aaa aaa ggg gaa gta gat gtt gtg tat tta gtt tat aac aaa ttt 888
254 Tyr Lys Lys Gly Glu Val Asp Val Val Tyr Leu Val Tyr Asn Lys Phe
255                160                165                170                175
257 att aac acg atg tcg caa aag cca gta ttg gaa aaa tta att cca tta 936
258 Ile Asn Thr Met Ser Gln Lys Pro Val Leu Glu Lys Leu Ile Pro Leu
259                180                185                190
261 cca gaa tta gat aat gat gaa tta ggc gaa aga aaa caa gtt tgg gat 984
262 Pro Glu Leu Asp Asn Asp Glu Leu Gly Glu Arg Lys Gln Val Trp Asp
263                195                200                205
265 tat att tac gaa cct gat gcg aaa gta tta tta gat aat tta ttg gtt 1032
266 Tyr Ile Tyr Glu Pro Asp Ala Lys Val Leu Leu Asp Asn Leu Leu Val
267                210                215                220
269 cgt tat tta gaa tct cag gtt tat caa gca gca gtt gaa aac ctt gct 1080
270 Arg Tyr Leu Glu Ser Gln Val Tyr Gln Ala Ala Val Glu Asn Leu Ala
271                225                230                235
273 tct gag caa gcc gct cga atg gtc gcc atg aaa gca gca aca gat aac 1128
274 Ser Glu Gln Ala Ala Arg Met Val Ala Met Lys Ala Ala Thr Asp Asn
275                240                245                250                255
277 gca ggt aac tta att aat gag tta cag tta gtc tat aac aaa gct cgt 1176
278 Ala Gly Asn Leu Ile Asn Glu Leu Gln Leu Val Tyr Asn Lys Ala Arg
279                260                265                270
281 caa gca agt att aca aat gaa tta aat gaa att gtt gcc ggt gca gca 1224

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Input Set : A:\Cpg.pto

Output Set: N:\CRF3\07192001\I545199C.raw

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282 Gln Ala Ser Ile Thr Asn Glu Leu Asn Glu Ile Val Ala Gly Ala Ala
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285 gca att taacaaatag aggatcggta atggcaactg gaaaaattgt acaaatcatc 1280
286 Ala Ile
288 ggtgcgggta ttgacgttga attcccacaa gatgcagtac caaaagtata tgatgcctta 1340
290 aatgttgaaa caggtttagt acttgaagtt caacaacaat taggtggtgg tgtagttcgc 1400
292 tgtatcgcaa tgggatcatc tgatggatta aaacgcgggt taagcgtaac aaatacgaat 1460
294 aacccaattt ctgttccagt gggaacgaaa acattgggtc gtatcatgaa cgtattgggt 1520
296 gaaccaatcg atgagcaagg tgaaatcggg gcagaagaga attggtctat tcaccgtgct 1580
298 ccaccaagtt atgaagaaca atctaacagt actgaacttt tagaaacggg aattaaagtt 1640
300 atcgacttag tttgtccggt tgcgaaaggg ggtaaagtag gtttattcgg tgggtcgggt 1700
302 gtcggtaaaa ccgtcaatat gatggaatta atccgtaaca tcgcaattga gcactcaggt 1760
304 tactctgtct ttgcgggggt aggtgagcgt acgcgtgaag gtaacgactt ctatcatgag 1820
306 atgaaagact ctaacgtatt agataaagt tctcttggtt atggtcaaat gaacgagcca 1880
308 ccaggttaacc gtttacgtgt ggcattaaca ggcttaacta tggcggaaaa attccgtgat 1940
310 gaaggtcgtg atgtcttatt cttcgttgat aa 1972
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323 20 25 30
325 Arg Lys Thr Gln Glu Arg Met Ser Ser Arg Pro Tyr Ser Glu Thr
326 35 40 45
328 Ile Arg Asn Val Ile Ser His Val Ser Lys Ala Thr Ile Gly Tyr Lys
329 50 55 60
331 His Pro Phe Leu Val Asp Arg Glu Val Lys Lys Val Gly Met Ile Val
332 65 70 75 80
334 Val Ser Thr Asp Arg Gly Leu Cys Gly Gly Leu Asn Val Asn Leu Phe
335 85 90 95
337 Lys Thr Val Leu Asn Glu Met Lys Glu Trp Lys Glu Lys Asp Val Ser
338 100 105 110
340 Val Gln Leu Ser Leu Ile Gly Ser Lys Ser Ile Asn Phe Phe Gln Ser
341 115 120 125
343 Leu Gly Ile Lys Ile Leu Thr Gln Asp Ser Gly Ile Gly Asp Thr Pro
344 130 135 140
346 Ser Val Glu Gln Leu Ile Gly Ser Val Asn Ser Met Ile Asp Ala Tyr
347 145 150 155 160
349 Lys Lys Gly Glu Val Asp Val Val Tyr Leu Val Tyr Asn Lys Phe Ile
350 165 170 175
352 Asn Thr Met Ser Gln Lys Pro Val Leu Glu Lys Leu Ile Pro Leu Pro
353 180 185 190
355 Glu Leu Asp Asn Asp Glu Leu Gly Glu Arg Lys Gln Val Trp Asp Tyr
356 195 200 205
358 Ile Tyr Glu Pro Asp Ala Lys Val Leu Leu Asp Asn Leu Leu Val Arg
359 210 215 220

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

F.Y.I. →
file://C:\Cr3\Outhoid\VSr1545199C.htm

7/19/01

VERIFICATION SUMMARY

DATE: 07/19/2001

PATENT APPLICATION: US/09/545,199C

TIME: 10:36:35

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\07192001\I545199C.raw

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L:3839 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:3840 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:3843 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:3844 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
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L:3848 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:3979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:3982 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:3985 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:5414 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:5415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:5426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:5427 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:5442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:5450 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
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L:11243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:90
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L:12304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:103